SEQUENCE LISTING

<110> Harper, Jeffrey W.
Elledge, Stephen J.

<120> F-BOX PROTEINS AND GENES

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<151> 1997-10-16

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gaaatatgg 129

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Leu Ser His Cys Glu Leu Ile Thr Asp Asp Gly Ile Leu His Leu Ser
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Arg Leu Ala Gln Thr Cys Lys Leu Leu Ser Gln His Cys Cys Asp Pro
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tttggataca ataaaaattt tatcctgata agatggcttg gtttgatagg aaatgccaga 1140

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acaatttttat agttgaactg taaaaaacta taaccattaa ttcttggtct acttgtaaga 1260
gtgagaattt acatgagctg cgctctctat ttttattaag gagagaagaa attaattcat 1320
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<210> 55
<211> 519
<212> PRT
<213> Homo sapiens

<400> 55
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Asp Glu Lys Ser Asp Lys Glu Ala Glu Val Ser Glu His Ser Thr Gly
20 25 30

Ile Thr His Leu Pro Pro Glu Val Met Leu Ser Ile Phe Ser Tyr Leu
35 40 45

Asn Pro Gln Glu Leu Cys Arg Cys Ser Gln Val Ser Met Lys Trp Ser
50 55 60

Gln Leu Thr Lys Thr Gly Ser Leu Trp Lys His Leu Tyr Pro Val His
65 70 75 80

Trp Ala Arg Gly Asp Trp Tyr Ser Gly Pro Ala Thr Glu Leu Asp Thr
85 90 95

Glu Pro Asp Asp Glu Trp Val Lys Asn Arg Lys Asp Glu Ser Arg Ala
100 105 110

Phe His Glu Trp Asp Glu Asp Ala Asp Ile Asp Glu Ser Glu Glu Ser
115 120 125

Ala Glu Glu Ser Ile Ala Ile Ser Ile Ala Gln Met Glu Lys Arg Leu
130 135 140

Leu His Gly Leu Ile His Asn Val Leu Pro Tyr Val Gly Thr Ser Val
145 150 155 160

Lys Thr Leu Val Leu Ala Tyr Ser Ser Ala Val Ser Ser Lys Met Val
165 170 175

Arg Gln Ile Leu Glu Leu Cys Pro Asn Leu Glu His Leu Asp Leu Thr
180 185 190

Gln Thr Asp Ile Ser Asp Ser Ala Phe Asp Ser Trp Ser Trp Leu Gly
195 200 205

Cys Cys Gln Ser Leu Arg His Leu Asp Leu Ser Gly Cys Glu Lys Ile
210 215 220

Thr Asp Val Ala Leu Glu Lys Ile Ser Arg Ala Leu Gly Ile Leu Thr
225 230 235 240

Ser His Gln Ser Gly Phe Leu Lys Thr Ser Thr Ser Lys Ile Thr Ser
245 250 255

Thr Ala Trp Lys Asn Lys Asp Ile Thr Met Gln Ser Thr Lys Gln Tyr
260 265 270

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Ala Cys Leu His Asp Leu Thr Asn Lys Gly Ile Gly Glu Glu Ile Asp
        275                 280                 285

Asn Glu His Pro Trp Thr Lys Pro Val Ser Ser Glu Asn Phe Thr Ser
    290                 295                 300

Pro Tyr Val Trp Met Leu Asp Ala Glu Asp Leu Ala Asp Ile Glu Asp
305                 310                 315                 320

Thr Val Glu Trp Arg His Arg Asn Val Glu Ser Leu Cys Val Met Glu
                325                 330                 335

Thr Ala Ser Asn Phe Ser Cys Ser Thr Ser Gly Cys Phe Ser Lys Asp
            340                 345                 350

Ile Val Gly Leu Arg Thr Ser Val Cys Trp Gln Gln His Cys Ala Ser
        355                 360                 365

Pro Ala Phe Ala Tyr Cys Gly His Ser Phe Cys Cys Thr Gly Thr Ala
    370                 375                 380

Leu Arg Thr Met Ser Ser Leu Pro Glu Ser Ser Ala Met Cys Arg Lys
385                 390                 395                 400

Ala Ala Arg Thr Arg Leu Pro Arg Gly Lys Asp Leu Ile Tyr Phe Gly
                405                 410                 415

Ser Glu Lys Ser Asp Gln Glu Thr Gly Arg Val Leu Leu Phe Leu Ser
            420                 425                 430

Leu Ser Gly Cys Tyr Gln Ile Thr Asp His Gly Leu Arg Val Leu Thr
        435                 440                 445

Leu Gly Gly Gly Leu Pro Tyr Leu Glu His Leu Asn Leu Ser Gly Cys
    450                 455                 460

Leu Thr Ile Thr Gly Ala Gly Leu Gln Asp Leu Val Ser Ala Cys Pro
465                 470                 475                 480

Ser Leu Asn Asp Glu Tyr Phe Tyr Tyr Cys Asp Asn Ile Asn Gly Pro
                485                 490                 495

His Ala Asp Thr Ala Ser Gly Cys Gln Asn Leu Gln Cys Gly Phe Arg
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Ala Cys Cys Arg Ser Gly Glu
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<210> 56
<211> 2276
<212> DNA
<213> Homo sapiens

<400> 56
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atgctgtcaa ttttcagcta tcttaatcct caagagttat gtcgatgcag tcaagtaagc 180
atgaaatggt ctcagctgac aaaaacggga tcgctttgga aacatcttta ccctgttcat 240
tgggccagag gtgactggta tagtggtccc gcaactgaac ttgatactga acctgatgat 300
gaatgggtga aaaataggaa agatgaaagt cgtgcttttc atgagtggga tgaagatgct 360
gacattgatg aatctgaaga gtctgcggag gaatcaattg ctatcagcat tgcacaaatg 420

gaaaaacgtt tactccatgg cttaattcat aacgttctac catatgttgg tacttctgta 480
aaaaccttag tattagcata cagctctgca gtttccagca aaatggttag gcagatttta 540
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aataaagaca ttaccatgca gtccaccaag cagtatgcct gtttgcacga tttaactaac 840
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cagaaggcca attcaaacaa acccacattt aaggttcttt aggattatag aataaattgg 2100
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<210> 57
<211> 39
<212> PRT
<213> Homo sapiens

<400> 57
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Thr Leu Leu Thr Cys Cys Leu Val Ser Lys Gln Trp Asn Lys Val Ile
20 25 30

Ser Ala Cys Thr Glu Val Trp
35

<210> 58
<211> 117
<212> DNA
<213> Homo sapiens

<400> 58
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tgctgcctcg tctctaaaca gtggaataag gtgataagtg cctgtacaga ggtgtgg 117

<210> 59
<211> 10
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 59
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<210> 60
<211> 21
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Synthetic

<400> 60
Lys Lys Glu Arg Leu Leu Asp Asp Arg His Asp Ser Gly Leu Asp Ser
1 5 10 15

Met Lys Asp Glu Glu
20